

payment of the fee therefor. Applicants respectfully request acceptance of the enclosed paper copy and computer readable form of the Sequence Listing. The Commissioner is hereby authorized to charge any additionally required fee, or credit any overpayment in fees, to Deposit Account No. 50-0320.

**AMENDMENT**

It is respectfully requested that the application be amended, without prejudice, without admission, without surrender of subject matter, and without any intention of creating any estoppel as follows:

**IN THE DRAWINGS:**

(12) Figures 1, 2 and 4 have been amended to include SEQ ID NO: 2-56, SEQ ID NO: 57-58 and SEQ ID NO: 139 respectively.

**IN THE SPECIFICATION:**

Page 10, line 21, please rewrite the paragraph thereat as follows:

Figure 2: Polynucleotide sequence of complete cDNA encoding human CTLA-4 and polypeptide sequence of the VLD of human CTLA-4. (SEQ ID NOS: 57-58)

Page 10, line 30, please rewrite the paragraph thereat as follows:

Figure 4: Schematic representation of the somatostatin polypeptide. Somatostatin (somatotropin release-inhibiting factor SRIF) is a cyclic 14-amino acid polypeptide. The cyclic nature is provided by a disulphide linkage between the cysteine residues at positions 3 and 14. The four residues which constitute the tip of the loop (Phe-Trp-Lys-Thr) are implicated in binding to members of the somatostatin receptor family. (SEQ ID NO: 139)

Page 27, line 1, please rewrite the table thereat as follows:

Table 4 CDR1 and CDR3 Inserts from a Representative Series of Library 2 Clones (SEQ ID NOS: 102-142)

CLONE	CDR1	CDR3
3M-2	ND1	LPSSTDTRAYS (SEQ ID NO: 102)
3M-3	QESGGRPG (SEQ ID NO: 103)	LPRGPPLLSL (SEQ ID NO: 104)
3M-5	SPGRCLN (SEQ ID NO: 105)	ND
3M-6	EWKREHGG (SEQ ID NO: 106)	LCPGACGCVY (SEQ ID NO: 107)
3M-7	NSGENEGG (SEQ ID NO: 108)	ND
3M-11	DKPVTKSG (SEQ ID NO: 109)	ND
3M-17	SPGACPE (SEQ ID NO: 110)	ND
3M-18	SPGKCDQ (SEQ ID NO: 111)	ND
3M-19	SPGMCAR (SEQ ID NO: 112)	LMYPPPYL (SEQ ID NO: 63)
3M-20	ND	PFLFLPCEFF (SEQ ID NO: 113)
3N-1	WTLGHHKLCEG (SEQ ID NO: 114)	LTFCLLALCS (SEQ ID NO: 115)
3N-2	SPGECYG (SEQ ID NO: 116)	SWLSTTXCLSSCS (SEQ ID NO: 117)
3N-3	SPGECQD (SEQ ID NO: 118)	LLGSLLSLSCFASLS (SEQ ID NO: 119)
3N-4	SPECQD (SEQ ID NO: 142)	SPGSLLSLSCFASXS (SEQ ID NO: 120)
3N-5	SPGRCTD (SEQ ID NO: 121)	VICHSSVCLSD/EVC (SEQ ID NOS: 122-123)
3N-6	ND	DLPSYLACSI (SEQ ID NO: 124)
3N-7	SPGRCTA (SEQ ID NO: 125)	ALCWDVIFYCSFPSY (SEQ ID NO: 126)
3N-8	ELFGHARYCKG (SEQ ID NO: 127)	VSITSPELCPVKVFD (SEQ ID NO: 128)
3N-9	SPGKVEN (SEQ ID NO: 129)	LFVPFVSP (SEQ ID NO: 130)
3N-12	SPGDLWV (SEQ ID NO: 131)	ESGLSPVSPCSLYSL (SEQ ID NO: 132)
3N-13	TSANGPYG (SEQ ID NO: 133)	PWAYRFLAVL (SEQ ID NO: 134)
3N-14	RKTREKYG (SEQ ID NO: 135)	ELMYPPPYLGI (SEQ ID NO: 136)
3N-15	SPGQELT (SEQ ID NO: 137)	ELFFLLYAPCYLFQR (SEQ ID NO: 138)

ND: Not Done

\*: UAG termination codon

Please replace the previously filed sequence listing with the enclosed papers titled --  
sequence listing.--